

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: March 1, 2001, 16:18:28 ; Search time 64.32 Seconds

(without alignments)
11.696 Million cell updates/sec

Title: US-09-331-631A-39

Perfect score: 54
Sequence: 1 CXXCXXXXXXXXXXCXXC 22

Scoring table: BLOSUM62PX
Gapop 10.0 , Gapext 0.5

Searched: 268485 seqs, 34193795 residues

Total number of hits satisfying chosen parameters: 268485

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	54	100.0	27	20	Y36499
2	54	100.0	31	21	Y70731
3	54	100.0	36	4	P30262
4	54	100.0	36	4	P30263
5	54	100.0	39	17	W05340
6	54	100.0	39	17	W05341
7	54	100.0	39	17	W05342
8	54	100.0	39	20	R07909
9	54	100.0	44	17	R38208
10	54	100.0	44	21	Y64770
11	54	100.0	57	21	Y57813
12	54	100.0	59	21	Y57812

13	54	100.0	62	21	Y57810	Human metallothion
14	54	100.0	63	21	Y57815	Sea urchin metallo
15	54	100.0	66	21	Y64780	Human 5' EST relat
16	54	100.0	70	21	Y75953	Murine skin cell p
17	54	100.0	72	19	W50389	Snake venom platelet
18	54	100.0	72	19	W50452	Snake venom platelet
19	54	100.0	72	19	W46214	Snake venom platelet
20	54	100.0	73	20	Y35935	Snake venom platelet
21	54	100.0	75	18	W21583	Extended human sec
22	54	100.0	76	20	Y02761	Alzheimer's diseas
23	54	100.0	76	21	Y68907	Human secreted pro
24	54	100.0	80	17	W05343	A mouse MDNM-2 pro
25	54	100.0	86	20	Y36458	Calisoga spider ve
26	54	100.0	92	21	Y69209	Fragment of human
27	54	100.0	93	20	Y36164	Amino acid sequenc
28	54	100.0	93	20	Y36211	Human secreted pro
29	54	100.0	100	21	Y65659	Human secreted pro
30	54	100.0	104	21	Y65661	C. elegans insulin
31	54	100.0	105	21	Y65660	C. elegans insulin
32	54	100.0	106	21	Y65655	C. elegans insulin
33	54	100.0	112	21	Y65658	C. elegans insulin
34	54	100.0	118	21	Y44985	Human epidermal pr
35	54	100.0	118	21	Y65662	C. elegans insulin
36	54	100.0	124	19	W56732	Nucleus-specific
37	54	100.0	125	12	R13329	HE4 epidermalys-spe
38	54	100.0	125	19	W81779	Human HE4 protein
39	54	100.0	128	21	Y44987	Human epidermal pr
40	54	100.0	149	8	P70057	Human insulin rece
41	54	100.0	150	8	P70058	Human epidermal gr
42	54	100.0	165	12	R10533	Prod. of pwc4812 u
43	54	100.0	169	20	Y60558	Human normal blad
44	54	100.0	170	20	Y29215	Amino acid sequenc
45	54	100.0	233	21	Y74791	Neisseria meningit

ALIGNMENTS

RESULT 1	
ID	Y36499 standard; Protein: 27 AA.
XX	
AC	Y36499;
XX	
DT	17-SEP-1999 (first entry)
XX	
DE	Fragment of human secreted protein encoded by gene 27.
XX	
KW	Human; secreted protein; cancer; tumour; developmental abnormality;
KW	foetal deficiency; blood disorder; immune system disorder; inflammation;
KW	autoimmune disease; allergy; Alzheimer's disease; cognitive disorder;
KW	schizophrenia; arthritis; asthma; psoriasis; sepsis; skin disorder;
KW	atherosclerosis; diabetes; cardiovascular disorder; kidney disorder;
KW	digestive disorder; endocrine disorder; infection; AIDS.
XX	
OS	Homo sapiens.
XX	
PN	W09931117-A1.
XX	
PD	24-JUN-1999.
XX	
PF	17-DEC-1998; 98WO-US27059.
XX	
PR	19-DEC-1997; 97US-0068369.
PR	18-DEC-1997; 97US-0068006.
PR	18-DEC-1997; 97US-0068007.
PR	18-DEC-1997; 97US-0068008.
PR	18-DEC-1997; 97US-0068053.
PR	18-DEC-1997; 97US-0068054.
PR	18-DEC-1997; 97US-0068057.
PR	18-DEC-1997; 97US-0068064.
PR	18-DEC-1997; 97US-0070923.
PR	19-DEC-1997; 97US-0068169.

PR 19-DEC-1997; 97US-0068365.
 PR 19-DEC-1997; 97US-0068367.
 PR 19-DEC-1997; 97US-0068368.
 XX
 PA (HUMA-) HUMAN GENOME SCI INC.
 XX
 PI Carter KC, Duan RD, Feng P, Ferrie AM, Florence C;
 PI Florence K, Greene JM, Janet F, Kyaw H, Moore PA;
 PI Ni J, Rosen CA, Ruben SM, Shi Y, Soppet DR, Wei Y;
 PI Yu G;
 XX
 DR WPI: 1999-418749/35.
 XX
 PT New isolated human genes encoding secreted polypeptides
 XX
 PS Disclosure: Page 466; 537pp; English.
 XX
 CC X97916 to X98029 represent 110 isolated human secreted protein genes.
 CC Y36224 to Y36727 represent the secreted proteins encoded by the 110
 CC human genes. The genes and their corresponding secreted polypeptides are
 CC useful for preventing, treating or ameliorating medical conditions,
 CC e.g. by protein or gene therapy. Also pathological conditions can be
 CC diagnosed by determining the amount of the new polypeptides in a sample
 CC or by determining the presence of mutations in the new genes. Specific
 CC uses are described for each of the 110 genes, based on which tissues they
 CC are most highly expressed in, and include developing products for the
 CC diagnosis or treatment of cancer, tumours, developmental abnormalities
 CC and foetal deficiencies, blood disorders, diseases of the immune system,
 CC autoimmune diseases, inflammation, allergies, Alzheimer's and cognitive
 CC disorders, schizophrenia, arthritis, asthma, psoriasis, sepsis, skin
 CC disorders, atherosclerosis, diabetes, cardiovascular disorders, kidney
 CC disorders, digestive/endocrine disorders, infections and AIDS. The
 CC polypeptides are also useful for identifying their binding partners.
 CC The sequences given in X97907 to X97915 and Y36223 are used in the
 CC exemplification of the present invention.
 XX
 SQ Sequence 27 AA:
 Query Match 100.0%; Score 54; DB 20; Length 27;
 Best Local Similarity 18.2%; Pred. No. 1e+02;
 Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;
 OY 1 CXXXCXXXXXXXXXXXXCXXC 22
 I:::|:::|:::|:::|:::|:::|
 Db 4 cpccclprgscrgcrafcsc 25
 RESULT 2
 ID Y70731 standard; protein: 31 AA.
 XX
 AC Y70731:
 XX
 DT 24-JUL-2000 (first entry)
 XX
 DE Wnt antagonist protein consensus sequence-1.
 XX
 KW Wnt antagonist; contraceptive; contraceptive vaccine; oocyte development;
 KW female primate contraception; oocyte viability.
 XX
 OS Synthetic.
 XX
 XX Key Location/Qualifiers
 FH Misc-difference 2
 FT /label= Unknown
 FT /note= "Xaa may be 9 amino acids in length; some
 FT amino acids may be absent."
 FT Misc-difference 4
 FT /label= Unknown
 FT /note= "Xaa may be 42 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 14

FT /label= Unknown
 FT Misc-difference 15
 FT /label= Unknown
 FT Misc-difference 16
 FT /label= Unknown
 FT Misc-difference 17
 FT /label= Unknown
 FT Misc-difference 18
 FT /label= Unknown
 FT Misc-difference 19
 FT /label= Unknown
 FT Misc-difference 21
 FT /label= Unknown
 FT /note= "Xaa may be 10 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 23
 FT /label= Unknown
 FT Misc-difference 24
 FT /label= Unknown
 FT Misc-difference 25
 FT /label= Unknown
 FT Misc-difference 27
 FT /label= Unknown
 FT /note= "Xaa may be 7 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 29
 FT /label= Unknown
 FT /note= "Xaa may be 27 amino acids in length; some
 FT amino acids may be absent"
 FT Misc-difference 31
 FT /label= Unknown
 FT /note= "Xaa may be 13 amino acids in length; some
 FT amino acids may be absent"
 XX
 PN WO200021555-A1.
 XX
 PD 20-APR-2000.
 XX
 PF 13-OCT-1999; 99WO-US23640.
 XX
 PR 15-OCT-1998; 98US-0104355.
 XX
 PA (HARD) HARVARD COLLEGE.
 XX
 PI McMahon AP, Parr BA, Vaino S;
 PI WPI: 2000-317845/27.
 DR
 XX
 PT Contraceptive composition for inhibiting oocyte development in a female
 PT primate comprises a Wnt polypeptide antagonist
 XX
 PS Claim 12; Page 44; 57pp; English.
 XX
 CC The patent discloses a method of female primate contraception comprising
 CC administering an antagonist of a Wnt polypeptide, inhibiting oocyte
 CC development. Wnt polypeptides are useful for promotive maturation of an
 CC immature oocyte. Wnt polypeptides are also useful for increasing the
 CC number of mature oocytes and to enhance oocyte viability. The present
 CC peptide is a consensus sequence of Wnt antagonist which inhibits the
 CC physiological activity of a Wnt polypeptide. Antagonistic polypeptides
 CC may contain a cysteine-rich domain.
 XX
 SQ Sequence 31 AA:
 Query Match 100.0%; Score 54; DB 21; Length 31;
 Best Local Similarity 63.6%; Pred. No. 1.1e+02;
 Matches 14; Conservative 8; Mismatches 0; Indels 0; Gaps 0;
 OY 1 CXXXCXXXXXXXXXXXXCXXC 22
 I:::|:::|:::|:::|:::|:::|
 Db 5 ccccccccxcccccxcccc 26

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RESULT 3
P30262 ID P30262 standard; peptide: 36 AA.
XX AC P30262:
XX DT 25-APR-1992 (first entry)
XX DE Sequence of peptide used to vaccinate against E. coli enterotoxin(s).
XX KW Vaccine; enterotoxin; diarrhoea; immunogen.
XX OS Escherichia coli.
XX FH Key Location/Qualifiers
FT MISC-difference 1..18 /label= Peptide P
XX PN EP93652-A.
XX PD 09-NOV-1983.
XX PF 26-APR-1983; 83EP-0072336.
XX PR 26-APR-1982; 82FR-0007179.
XX PA (INSP ) INST PASTEUR.
XX PI (CNRS ) CENT NAT RECH SCT.
XX PI Tartar A, Duflot E, Boquet P;
XX DR WPI; 1983-816301/46.
XX PT Peptide(s) used to vaccinate against E. coli enterotoxin(s) -
PT contg. e.g. asparagine threonine phenylalanine tyrosine cysteine
PT cysteine glutamic acid leucine cysteine asparagine
XX PT sequences
XX PS Claim 1; Page 40; 50pp; French.
XX CC The inventors claim peptides of formula (P)n (see FT; see also
CC P30263) having 4n-18n amino acids and pref. being laevorotatory
CC (where n is 1 or 2). In P30262 and P30263, N=2. When n is 2, the
CC peptide comprises two peptide sequences Pi which may be the same or
CC different, each having 4-18 amino acids chosen from the peptide P SQ
CC in P30262 or P30263. The two P sequences may be joined (a) by a
CC disulphide bond or (b) by a bond formed between a carboxyl gp. of
CC one sequence of an amino gp. of the other.
XX SQ Sequence 36 AA:

Query Match 100.0%; Score 54; DB 4; Length 36;
Best Local Similarity 18.2%; Pred. No. 1.3e+02;
Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

QY 1 CXXXXXXXXXXXXCXXC 22
|::|::|::|::|::|::|::|
Db 6 celcnpacagcnytfccelc 27

RESULT 4
P30263 ID P30263 standard; peptide: 36 AA.
XX AC P30263:
XX DT 25-APR-1992 (first entry)
XX DE Sequence of peptide used to vaccinate against E. coli enterotoxin(s).
XX KW Vaccine; enterotoxin; diarrhoea; immunogen.

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XX OS Escherichia coli.
XX FH Key Location/Qualifiers
FT MISC-difference 1..18 /label= Peptide P
XX PN EP93652-A.
XX PD 09-NOV-1983.
XX PF 26-APR-1983; 83EP-0072336.
XX PR 26-APR-1982; 82FR-0007179.
XX PA (INSP ) INST PASTEUR.
XX PI (CNRS ) CENT NAT RECH SCT.
XX PI Tartar A, Duflot E, Boquet P;
XX DR WPI; 1983-816301/46.
XX PT Peptide(s) used to vaccinate against E. coli enterotoxin(s) -
PT contg. e.g. asparagine threonine phenylalanine tyrosine cysteine
PT cysteine glutamic acid leucine cysteine asparagine
XX PT sequences
XX PS Claim 1; Page 40; 50pp; French.
XX CC The inventors claim peptides of formula (P)n (see FT; see also
CC P30263) having 4n-18n amino acids and pref. being laevorotatory
CC (where n is 1 or 2). In P30262 and P30263, N=2. When n is 2, the
CC peptide comprises two peptide sequences Pi which may be the same or
CC different, each having 4-18 amino acids chosen from the peptide P SQ
CC in P30262 or P30263. The two P sequences may be joined (a) by a
CC disulphide bond or (b) by a bond formed between a carboxyl gp. of
CC one sequence of an amino gp. of the other.
XX SQ Sequence 36 AA:

Query Match 100.0%; Score 54; DB 4; Length 36;
Best Local Similarity 18.2%; Pred. No. 1.3e+02;
Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

QY 1 CXXXXXXXXXXXXCXXC 22
|::|::|::|::|::|::|::|
Db 6 celcypacagcnytfccelc 27

RESULT 5
W05340 ID W05340 standard; peptide: 39 AA.
XX AC W05340:
XX DT 15-APR-1997 (first entry)
XX DE Callisoga spider venom peptide A, used as insecticide.
XX KW Callisoga: spider; venom; insecticide; recombinant; baculovirus; pest;
XX KW tobacco budworm; Heliothis virescens; low mammalian toxicity.
XX OS Callisoga sp.
XX PN W09625041-A1.
XX PD 22-AUG-1996.
XX PF 16-FEB-1996; 96WO-US02030.
XX PR 17-FEB-1995; 95US-0390882.
XX

```

PA (NPSp-) NPS PHARM INC.
 XX Johnson JH, Kral RM, Krapcho K;
 XX
 DR WPI: 1996-393030/39.
 DR N-PSDB: T39769.
 XX
 PT New insecticidal peptide(s) from venom of *Calisoga* spider - having
 PT low toxicity to mammals, useful for controlling insect pests
 PS
 PS Claim 2: Page 33; 53pp; English.
 XX
 CC W05340 is a peptide derived from the venom of spiders of the
 CC genus *Calisoga*. It is useful as an insecticide, having a neurotoxic
 CC effect on *Heliothis virescens* (tobacco budworm). The peptide is
 CC derived from a larger protein sequence including a 41 amino acid (aa)
 CC signal peptide region (see W05343) and is preferably recombinantly
 CC administered to insects using a baculovirus host expression system
 CC (or other natural insect pathogen, e.g. *Bacillus*). The peptide has
 CC a low toxicity to mammals and can be recombinantly produced on a
 CC large scale.
 CC
 SQ Sequence 39 AA:

Query Match 100.0%; Score 54; DB 17; Length 39;
 Best Local Similarity 18.2%; Pred. No. 1.4e+02;
 Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXXXXXXXXXXXXXXXXC 22
 ID W05341 standard; peptide: 39 AA.
 DB 15 csqncgtwctcylrkdkpskcc 36

RESULT 6
 W05341
 ID W05341 standard; peptide: 39 AA.
 AC W05341:
 DE 15-APR-1997 (first entry)
 DE *Calisoga* spider venom peptide B, used as insecticide.
 KW *Calisoga*; spider; venom; insecticide; recombinant; baculovirus; pest;
 KW tobacco budworm; *Heliothis virescens*; low mammalian toxicity.
 OS *Calisoga* sp.
 XX W09625041-A1.
 XX
 XX 22-AUG-1996.
 XX 16-FEB-1996; 96WO-US02030.
 XX 17-FEB-1995; 95US-0390882.
 XX (NPSp-) NPS PHARM INC.
 XX Johnson JH, Kral RM, Krapcho K;
 XX
 DR WPI: 1996-393030/39.
 DR N-PSDB: T39770
 XX
 PT New insecticidal peptide(s) from venom of *Calisoga* spider - having
 PT low toxicity to mammals, useful for controlling insect pests
 PS
 PS Claim 3: Page 39; 53pp; English.
 XX
 CC W05341 is a peptide derived from the venom of spiders of the
 CC genus *Calisoga*. It is useful as insecticides having a neurotoxic
 CC effect on *Heliothis virescens* (tobacco budworm). The peptide is
 CC preferably recombinantly administered to insects using a baculovirus

CC host expression system (or other natural insect pathogen, e.g.
 CC *Bacillus*). The peptide has a low toxicity to mammals and can be
 CC recombinantly produced on a large scale.
 CC
 SQ Sequence 39 AA:

Query Match 100.0%; Score 54; DB 17; Length 39;
 Best Local Similarity 18.2%; Pred. No. 1.4e+02;
 Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXXXXXXXXXXXXXXXXC 22
 ID W05342 standard; peptide: 39 AA.
 DB 15 csqncgtwctcylrkdkpskcc 36

RESULT 7
 W05342
 ID W05342 standard; peptide: 39 AA.
 AC W05342:
 DE 15-APR-1997 (first entry)
 DE *Calisoga* spider venom peptide C, used as insecticide.
 KW *Calisoga*; spider; venom; insecticide; recombinant; baculovirus; pest;
 KW tobacco budworm; *Heliothis virescens*; low mammalian toxicity.
 OS *Calisoga* sp.
 XX W09625041-A1.
 XX
 XX 22-AUG-1996.
 XX 16-FEB-1996; 96WO-US02030.
 XX 17-FEB-1995; 95US-0390882.
 XX (NPSp-) NPS PHARM INC.
 XX Johnson JH, Kral RM, Krapcho K;
 XX
 DR WPI: 1996-393030/39.
 DR N-PSDB: T39770.
 XX
 PT New insecticidal peptide(s) from venom of *Calisoga* spider - having
 PT low toxicity to mammals, useful for controlling insect pests
 PS
 PS Claim 4: Page 40; 53pp; English.
 XX
 CC W05342 is a peptide derived from the venom of spiders of the
 CC genus *Calisoga*. It is useful as insecticides having a neurotoxic
 CC effect on *Heliothis virescens* (tobacco budworm). The peptide is
 CC derived from a larger protein sequence including a 41 amino acid (aa)
 CC signal peptide region (see W05343) and is preferably recombinantly
 CC administered to insects using a baculovirus host expression system
 CC (or other natural insect pathogen, e.g. *Bacillus*). The peptide has
 CC a low toxicity to mammals and can be recombinantly produced on a
 CC large scale.
 CC
 SQ Sequence 39 AA:

Query Match 100.0%; Score 54; DB 17; Length 39;
 Best Local Similarity 18.2%; Pred. No. 1.4e+02;
 Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXXXXXXXXXXXXXXXXC 22
 ID W05341 standard; peptide: 39 AA.
 DB 15 csqncgtwctcylrkdkpskcc 36

RESULT	8
ID	Y07909
AC	Y07909 standard; Protein; 39 AA.
XX	
XX	Y07909;
DT	06-JUL-1999 (first entry)
DE	
XX	Human secreted protein fragment encoded from gene 58.
XX	
KW	Human; secreted protein; treatment; prevention; protein therapy; AIDS;
KW	gene therapy; diagnosis; cancer; tumour; neurodegenerative disorder;
KW	developmental abnormality; fetal deficiency; blood disorder; leukemia;
KW	immune system disease; autoimmune disease; hepatic disease; lymphoma;
KW	renal disease; inflammation; allergy; Alzheimer's disease; schizophrenia
KW	cognitive disorder; prostate disease; skeletal; cardiac; muscle disorder
KW	pulmonary disorder; transplant rejection; osteoclast; osteoporosis;
KW	arthritis; malignancy; digestive; endocrine; infection.
OS	
XX	Homo sapiens.
PN	
PD	MO9918208-A1.
XX	
XX	15-APR-1999.
PF	
PE	01-OCT-1998; 98MO-US020775.
XX	
PR	02-OCT-1997; 97US-0060884.
PR	02-OCT-1997; 97US-0060833.
PR	02-OCT-1997; 97US-0060836.
PR	02-OCT-1997; 97US-0060837.
PR	02-OCT-1997; 97US-0060838.
PR	02-OCT-1997; 97US-0060839.
PR	02-OCT-1997; 97US-0060843.
PR	02-OCT-1997; 97US-0060862.
PR	02-OCT-1997; 97US-0060866.
PR	02-OCT-1997; 97US-0060874.
XX	
PA	(HUMA-) HUMAN GENOME SCI INC.
PI	
PI	Carter KC, Duan DR, Endress GA, Feng P, Ferrie AM;
PI	Florence KA, Greene JM, Janat F, Lafleur DW, Ni J;
PI	Rosen CA, Ruben SM, Shi Y, Young P, Yu G;
DR	WIPI: 1999-264022/22.
N-PSDB:	XJ37508.
XX	
XX	New isolated human genes and the secreted polypeptides they encode
PT	
XX	Claim 1b; Page 306; 368pp; English.
PS	
XX	
CC	This invention describes novel isolated human genes and the secreted
CC	proteins they encode. The products of the invention are useful for
CC	preventing, treating or ameliorating medical conditions, e.g. by protein
CC	or gene therapy. Also pathological conditions can be diagnosed by
CC	determining the amount of the new polypeptides in a sample or by
CC	determining the presence of mutations in the new polynucleotides.
CC	Specific uses are described for each of the 101 polynucleotides, based on
CC	which tissues they are most highly expressed in, and include developing
CC	products for the diagnosis or treatment of cancer, tumours,
CC	neurodegenerative disorders, developmental abnormalities and fetal
CC	deficiencies, blood disorders, leukemias, diseases of the immune system,
CC	autoimmune diseases, hepatic and renal disease, lymphomas, inflammation,
CC	allergies, Alzheimer's and cognitive disorders, schizophrenia, prostate
CC	disease, skeletal or cardiac muscle disorders, pulmonary disorders,
CC	transplant rejection, disorders involving osteoclasts such as
CC	osteoporosis, arthritis or malignancies, digestive/endocrine disorders,
CC	infections and AIDS. The human secreted proteins of the invention are
CC	represented in XJ37451-XJ37552.
XX	
Sequence	39 AA;
50	

```

Query Match          100.0%; Score 54; DB 20; length 39;
Best Local Similarity 18.2%; Pred. No. 1.4e+02;
Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;
Qy 1 CXXXXXXXXXXXXXXXXCXXC 22
|::|::|::|::|::|::|::|::|
Db 6 cfracemcs1sg11n1c1qsc 27

```

RESULT 9

R98208
ID R98208 standard; Protein; 44 AA.
YV

AC R98208;

DT 30-DEC-1996 (first entry)
 YY

DE Nucleotide used in production of MSH/MOMULV chimeric sequence.
XY

KW 10A1 murine leukaemia virus: N2B-9-1 murine leukaemia virus.
KW Moloney murine leukaemia virus; gp70; 4070A retrovirus; retrovirus.

KW single chain antibody; envelope protein; ss.

05 Synthetic.

PN W09630504-A1.

PD 03-OCT-1996.

PF 22-MAR-1996; 96WQ-US03908.
XX

PR	24-MAR-1993;	9305-0409648.
XX		

PA (UYSC-) UNIV SOUTHERN CALIFORNIA.

PI Anderson W, Chiang YL, Januszewski M, Mackrell AJ;

XX
DD
WDT 1006 AEECEC /AE

XX	Call-targeted retroviral vector particles having unique restriction
PT	

PI modified with targeting polypeptide

Example 2, page 30, 199p, English, XX

CC therapy to deliver a heterologous gene to a target cell for

targeted retroviral vector particles comprise an envelope protein

CC antibody), or in the case of moloney murine leukaemia virus (MolMuLV) alpha interferon stimulates bone marrow (BM) mu-

oligonucleotides (R98207, R98208) were used to substitute sequences

residues 600-700 of MOMLV envelope protein (see W04248).

Sequence 44 AA;

Query Match	100.08; Score 54; DB 17; Length 44;
-------------	-------------------------------------

Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps

QY	1	CXXXXXXXXXXXXXXXXC	22
		[...]	

Db 15 caaqccqtaataacctccctc 36

RESULT 10

Y64770

ID Y64770 standard; Protein; 44 AA.
 AC Y64770;
 DT 01-FEB-2000 (first entry)
 DE Human 5' EST related polypeptide seq ID NO:931.
 XX
 XX Human; 5' EST: expressed sequence tag; secreted protein; diagnosis;
 KW gene therapy; chromosome mapping; upstream regulatory sequence;
 KW forensic; location; development; protein synthesis; stability;
 KW regulation; identification.
 XX
 OS Homo sapiens.
 PN W0953051-A2.
 XX
 XX 21-OCT-1999.
 PD
 XX
 XX 09-APR-1999; 99WO-IB00712.
 PF
 XX 09-APR-1998; 98US-0057719.
 PR 28-APR-1998; 98US-0069047.
 XX
 XX (GEST) GENSET.
 PA
 P1 Dunas Milne Edwards J, Duclert A, Giordano J;
 DR WPI: 2000-038446/03.
 DR N-PSDB: 242384.
 PT Novel secreted protein 5' expressed sequence tag sequences used in
 PT diagnostic, forensic, gene therapy, and chromosome mapping procedures
 XX
 PS Claim 3: Page 637; 837pp; English.
 XX
 XX 242265 to 243075 represent novel 5' expressed sequence tag (EST)
 CC sequences, corresponding to human secreted proteins. Y64651 to Y64438
 CC represent the EST-related proteins corresponding to 242265 to 243052.
 CC The 5' ESTs can be used for producing secreted human gene products.
 CC They can be used to identify and isolate 5' untranslated regions (UTRs)
 CC and upstream regulatory regions which control the location, development
 CC stage, rate, and quantity of protein synthesis, as well as stability of
 CC mRNA. The ESTs are also useful as probes for chromosome mapping, and to
 CC obtain full length cDNA clones. The ESTs can also be used in forensic
 CC procedures to identify individuals, or in diagnostic procedures to
 CC identify individuals having genetic diseases resulting from abnormal
 CC gene expression. The products may also be used in gene therapy protocols.
 CC The nucleic acids encoding signal peptides can be used for directing
 CC extracellular secretion of a polypeptide or the insertion of a
 CC polypeptide into a membrane, or importing a polypeptide into a cell.
 CC The proteins encoded by the EST sequences may be useful in treating a
 CC variety of human conditions. Secreted proteins have therapeutic value,
 CC and the identification of new secreted proteins is valuable. 242249 to
 CC 242264 and Y64664 to Y64650 represent sequences used in the
 CC exemplification of the present invention.
 XX
 XX Sequence 44 AA:

```

Query Match      100.0%; Score 54; DB 21; Length 44;
Best Local Similarity 18.2%; Pred. No. 1.5e+02;
Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

QY      1 CXXXCXXXXXXXXXXXXCXXC 22
        |:::|:::|:::|:::|:::|
Db       15 CVCVCVCVCVCPYGMYLVCVCVC 36

RESULT 11
Y57813
ID      Y57813 standard; protein; 57 AA.
XX

```

AC Y57813;
XX
DT 22-MAR-2000 (first entry)
XX
DE Crab metallothionein Class I amino acid sequence.
XX
KW Metallothionein; metal recovery; remediation; heavy metal;
XX precious metal; phytochelatin; green algae; Chlamydomonas reinhardtii.
OS Eubrachyura.
XX
PN WO9960838-A1.
XX
PD 02-DEC-1999.
XX
PF 28-MAY-1999; 99WO-US12007.
XX
PR 28-MAY-1998; 98US-0087374.
XX
PA (OHIS) UNIV OHIO STATE RES FOUND.
XX
PI Sayre RT, Trajna SJ;
XX
DR WPI: 2000-086646/07.
XX
PT Novel method for metal recovery, remediation and separation -
XX
PS Disclosure; Page 6; 86pp; English.
XX
CC The present invention describes a transgenic algal cell (I) of the
XX genus Chlamydomonas comprising reproductive genetic material comprising
XX a nucleotide sequence capable of expressing chicken type I
XX Metallothionein. Also described is a method of removing metal from
XX an aqueous medium containing at least one dissolved or suspended
XX metal. The transgenic algae are used for the selective separation of
XX metals, particularly the separation of precious and desirable metals
XX such as gold and uranium, from other metals such as cadmium, zinc and
XX copper. The method can be used to facilitate the selective recovery of
XX precious and rare metals from mineral sources where aqueous media can
XX be used, such as in natural surface water flows, ground water and where
XX water may be introduced. The method is suitable for well-drilling,
XX soil and water remediation arts, mining fields, and industrial
XX engineering. The present sequence represents a Class I metallothionein
XX given in the present invention.
XX
Sequence 57 AA:

```

Query Match      100.0%; Score 54; DB 21; Length 57;
Best Local Similarity 18.2%; Pred. NO. 1.9e+02;
Matches 4; Conservative 18; Mismatches 0; Indels 0; Gaps 0;

Qy      1 CXXXCXXXXXXXXXXCXCXC 22
      |:::|:::|:::|:::|:::|
Db      16 cktgckctscrcpocqcsyc 37

RESULT 12
y57812
ID      y57812 standard; protein; 59 AA.
XX
XX      AC      y57812;
XX
XX      DT      22-MAR-2000 (first entry)
XX
DE      Trout metallothionein Class I amino acid sequence.
XX
XX
XX      DE      Metallothionein; metal recovery; remediation; heavy metal;
XX      KW      precious metal; phytochelatin; green algae; Chlamydomonas reinhardtii.
XX      XX
OS      Salmo sp.
XX
XX      W09960838-A1.
XX      PN

```


50 Sequence 63 AA;

qy	1	CAXXCXXXXXXXXXXCXXC	22
		:: :: :: :: :: :: ::	
Db	34	ccgcinaackcangckcgsgc	55

ID	Y64780	standard; Protein; 66 AA.
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AC Y64780;

DT 01-FEB-2000 (first entry)

Human 5' EST related polypeptide SEQ ID NO:941.

KM Hunnan, 5' ESR; expressed sequence tag; secreted protein; diagnosis
KM gene therapy; chromosome mapping; upstream regulatory sequence;
KM forensic; location; development; protein synthesis; stability;
KM regulation; identification.

OS Homo sapiens.

PN W099953051-A2

PD 21-OCT-1999

PF 09-APR-1999; 99WO-IB00712
VY

PR	09-APR-1998;	9805-0057719
DE	38-APR-1998;	9805-0050047

XX
PA (CECT) CENSEPT

PI Dumas Milne Edwards J, Duclert A, Giordano J;
v

DR WPI; 2000-038446/03.
DR N-DCDD; 743304

DR N-PSDB; Z42394.

PT Novel secreted protein 5' expressed sequence tag sequences used in
diagnostic, forensic, gene therapy, and chromosome mapping procedures
XX
PS Claim 3; Page 640; 837pp; English.

Claim 3; Page 640; 837pp; English

CC 242265 243075 represent novel5' expressed sequence tag (EST) sequences, corresponding to human secreted proteins. Y64651 to Y65438 CC represent the EST-related proteins corresponding to 242265 to 2423052. CC The 5' ESTs can be used for producing secreted human gene products. CC They can be used to identify and isolate 5' untranslated regions (UTRs) CC and upstream regulatory regions which control the location, development CC stage, rate, and quantity of protein synthesis, as well as stability of CC mRNA. The ESTs are also useful as probes for chromosome mapping, and to CC obtain full length cDNA clones. The ESTs can also be used in forensic CC procedures to identify individuals, or in diagnostic procedures to CC identify individuals having genetic diseases resulting from abnormal CC gene expression. The products may also be used in gene therapy protocols CC The nucleic acids encoding signal peptides can be used for directing CC extracellular secretion of a polypeptide or the insertion of a

CC polypeptide into a membrane, or importing a polypeptide into a cell.
CC The proteins encoded by the EST sequences may be useful in treating a
CC variety of human conditions. Secreted proteins have therapeutic value.
CC and the identification of new secreted proteins is valuable. 242249 ld
CC 242264 and Y64644 to Y64650 represent sequences used in the
CC exemplification of the present invention.

SQ Sequence 66 AA;

Query Match	100.0%;	Score 54;	DB 21;	Length 66;
Best Local Similarity	18.2%;	Pred. NO. 2.2e+02;		
Matches	4;	Conservative	18;	Mismatches 0;
			Indels	0;
			Gaps	0;

```
QY 1 CXXXXCXXXXXXXXXXXXCXXXC 22
    |::|:::|:::|:::|:::|
Db 17 clvscvlcvpcpvcwmccvcwc 38
```

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Search completed: March 1, 2001, 16:18:29
Job time: 498 sec
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